
Sequence Listing could not be accepted due to errors.

See attached Validation Report.

If you need help call the Patent Electronic Business Center at (866) 217-9197 (toll free).

Reviewer: Durreshwar Anjum

Timestamp: [year=2009; month=12; day=24; hr=13; min=50; sec=43; ms=621;

Reviewer Comments:

Seq Id # 6

Leu Ala Arg Ala Val Phe Ala Ser Ser Arg Thr Phe Gly Pro Ala Ser 995 1000 1005

Phe Ala Arg Ile Glu Thr Ala Phe Ala Asn Leu Tyr Pro Gly Glu 1010 1015 1020

Gln Pro Leu Cys Leu Cys Arg Gly Gly Asn Val Ala Tyr Thr Val 1025 1030 1035

Please move a space between the bases Gly Pro 1005

Please make necessary changes.

Validated By CRFValidator v 1.0.3

Application No: 10591787 Version No: 2.0

Input Set:

Output Set:

Started: 2009-12-03 19:02:14.043 **Finished:** 2009-12-03 19:02:26.589

Elapsed: 0 hr(s) 0 min(s) 12 sec(s) 546 ms

Total Warnings: 5
Total Errors: 1

No. of SeqIDs Defined: 91

Actual SeqID Count: 91

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E	323	Invalid/missing amino acid numbering SEQID (6) POS (1	.006)
W	213	Artificial or Unknown found in <213> in SEQ ID (7)	

SEQUENCE LISTING

<110>	Nagaike, Kazuhiro
	Mori, Yasuko
	Gomi, Yasuyuki
	Takahashi, Michiaki
	Kouichi, Yamanishi
<120>	Recombinant varicella-zoster virus
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                                                    Met Asp Thr
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                                                                     285
Pro Pro Met Gln Arg Ser Thr Pro Gln Arg Ala Gly Ser Pro Asp Thr
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5

10

15

_	gag Glu		_	_	_	_	_		-	_		_	_	_		333
	gcc Ala			-		_	-	_		-	-					381
	aac Asn		_	_				-				-				429
	gta Val	_					-		-	_	-	-	-		-	477
	cta Leu 85			-	_		_	_	, ,	_		_	-		1 2	525
	tcg Ser		-		-	-	-		-				-		-	573
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-	tct Ser			_	_		-						_	-		669
_	ggg	-	_		-		_			_		_	_	-		717
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Gln 180	cgg Arg	Glu	Asp	Gln	Arg 185	Ser	Ile	Pro	Arg	Gln 190	Thr	Thr	Arg	Gly	Asn 195	813
Ser	cca Pro	Gly	Leu	Pro 200	Ser	Val	Val	His	Arg 205	Asp	Arg	Gln	Thr	Gln 210	Ser	861
Ile	tcg Ser	Gly	Lys 215	Lys	Pro	Gly	Asp	Glu 220	Gln	Ala	Gly	His	Ala 225	His	Ala	909
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-	-	tta Leu	_		_	_	_	_	_		_		-		-	2	301
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700 705

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_	_			_	gga Gly 745	_	-							_		2493
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_	-	_	_	cgc Arg 1030				_	gca Ala 1035	Tyr	acc Thr		_		3348
_	gcg Ala			aag Lys 1045		_	-		ctg Leu 1050	Ser	ccc Pro	_	-		3393
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				ccc Pro 1150		_			_			ctg Leu			3708
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